

### AMENDMENTS TO THE CLAIMS

1. **(Currently amended)** An isolated polypeptide having at least 80% amino acid sequence identity to:
  - (a) the amino acid sequence of the polypeptide of SEQ ID NO:90 ~~shown in FIG. 90 (SEQ ID NO:90)~~;
  - (b) the amino acid sequence of the polypeptide of SEQ ID NO:90 ~~shown in FIG. 90 (SEQ ID NO:90)~~, lacking its associated signal peptide;
  - (c) the amino acid sequence of the extracellular domain of the polypeptide of SEQ ID NO:90, wherein the extracellular domain is amino acids 29-50 or 125-140 ~~shown in FIG. 90 (SEQ ID NO:90)~~;
  - (d) the amino acid sequence of the extracellular domain of the polypeptide of SEQ ID NO:90, wherein the extracellular domain is amino acids 29-50 or 125-140 ~~shown in FIG. 90 (SEQ ID NO:90)~~, lacking its associated signal peptide; or
  - (e) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209399;  
wherein said isolated polypeptide is more highly expressed in kidney tumor compared to normal kidney tissue, or wherein said isolated polypeptide is encoded by a polynucleotide that is more highly expressed in kidney tumor compared to normal kidney tissue.
2. **(Currently amended)** The isolated polypeptide of claim 1 having at least 85% amino acid sequence identity to:
  - (a) the amino acid sequence of the polypeptide of SEQ ID NO:90 ~~shown in FIG. 90 (SEQ ID NO:90)~~;
  - (b) the amino acid sequence of the polypeptide of SEQ ID NO:90 ~~shown in FIG. 90 (SEQ ID NO:90)~~, lacking its associated signal peptide;
  - (c) the amino acid sequence of the extracellular domain of the polypeptide of SEQ ID NO:90, wherein the extracellular domain is amino acids 29-50 or 125-140 ~~shown in FIG. 90 (SEQ ID NO:90)~~;

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(d) the amino acid sequence of the extracellular domain of the polypeptide of SEQ ID NO:90, wherein the extracellular domain is amino acids 29-50 or 125-140~~shown in FIG. 90 (SEQ ID NO:90)~~, lacking its associated signal peptide; or

(e) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209399;  
wherein said isolated polypeptide is more highly expressed in kidney tumor compared to normal kidney tissue, or wherein said isolated polypeptide is encoded by a polynucleotide that is more highly expressed in kidney tumor compared to normal kidney tissue.

3. **(Currently amended)** The isolated polypeptide of claim 1 having at least 90% amino acid sequence identity to:

(a) the amino acid sequence of the polypeptide of SEQ ID NO:90 ~~shown in FIG. 90 (SEQ ID NO:90)~~;

(b) the amino acid sequence of the polypeptide of SEQ ID NO:90 ~~shown in FIG. 90 (SEQ ID NO:90)~~, lacking its associated signal peptide;

(c) the amino acid sequence of the extracellular domain of the polypeptide of SEQ ID NO:90, wherein the extracellular domain is amino acids 29-50 or 125-140~~shown in FIG. 90 (SEQ ID NO:90)~~;

(d) the amino acid sequence of the extracellular domain of the polypeptide of SEQ ID NO:90, wherein the extracellular domain is amino acids 29-50 or 125-140~~shown in FIG. 90 (SEQ ID NO:90)~~, lacking its associated signal peptide; or

(e) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209399;  
wherein said isolated polypeptide is more highly expressed in kidney tumor compared to normal kidney tissue, or wherein said isolated polypeptide is encoded by a polynucleotide that is more highly expressed in kidney tumor compared to normal kidney tissue.

4. **(Currently amended)** The isolated polypeptide of claim 1 having at least 95% amino acid sequence identity to:

(a) the amino acid sequence of the polypeptide of SEQ ID NO:90 ~~shown in FIG. 90 (SEQ ID NO:90)~~;

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- (b) the amino acid sequence of the polypeptide of SEQ ID NO:90 ~~shown in FIG. 90 (SEQ ID NO:90)~~, lacking its associated signal peptide;
- (c) the amino acid sequence of the extracellular domain of the polypeptide of SEQ ID NO:90, wherein the extracellular domain is amino acids 29-50 or 125-140 ~~shown in FIG. 90 (SEQ ID NO:90)~~;
- (d) the amino acid sequence of the extracellular domain of the polypeptide of SEQ ID NO:90, wherein the extracellular domain is amino acids 29-50 or 125-140 ~~shown in FIG. 90 (SEQ ID NO:90)~~, lacking its associated signal peptide; or
- (e) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209399;  
wherein said isolated polypeptide is more highly expressed in kidney tumor compared to normal kidney tissue, or wherein said isolated polypeptide is encoded by a polynucleotide that is more highly expressed in kidney tumor compared to normal kidney tissue.

5. **(Currently amended)** The isolated polypeptide of claim 1 having at least 99% amino acid sequence identity to:

- (a) the amino acid sequence of the polypeptide of SEQ ID NO:90 ~~shown in FIG. 90 (SEQ ID NO:90)~~;
- (b) the amino acid sequence of the polypeptide of SEQ ID NO:90 ~~shown in FIG. 90 (SEQ ID NO:90)~~, lacking its associated signal peptide;
- (c) the amino acid sequence of the extracellular domain of the polypeptide of SEQ ID NO:90, wherein the extracellular domain is amino acids 29-50 or 125-140 ~~shown in FIG. 90 (SEQ ID NO:90)~~;
- (d) the amino acid sequence of the extracellular domain of the polypeptide of SEQ ID NO:90, wherein the extracellular domain is amino acids 29-50 or 125-140 ~~shown in FIG. 90 (SEQ ID NO:90)~~, lacking its associated signal peptide; or
- (e) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209399;  
wherein said isolated polypeptide is more highly expressed in kidney tumor compared to normal kidney tissue, or wherein said isolated polypeptide is encoded by a polynucleotide that is more highly expressed in kidney tumor compared to normal kidney tissue.

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6. **(Currently amended)** An isolated polypeptide comprising:
- (a) the amino acid sequence of the polypeptide of SEQ ID NO:90 ~~shown in FIG. 90 (SEQ ID NO:90)~~;
  - (b) the amino acid sequence of the polypeptide of SEQ ID NO:90 ~~shown in FIG. 90 (SEQ ID NO:90)~~, lacking its associated signal peptide;
  - (c) the amino acid sequence of the extracellular domain of the polypeptide of SEQ ID NO:90, wherein the extracellular domain is amino acids 29-50 or 125-140 ~~shown in FIG. 90 (SEQ ID NO:90)~~;
  - (d) the amino acid sequence of the extracellular domain of the polypeptide of SEQ ID NO:90, wherein the extracellular domain is amino acids 29-50 or 125-140 ~~shown in FIG. 90 (SEQ ID NO:90)~~, lacking its associated signal peptide; or
  - (e) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209399.
7. **(Currently amended)** The isolated polypeptide of claim 6 comprising the amino acid sequence of the polypeptide of SEQ ID NO:90 ~~shown in FIG. 90 (SEQ ID NO:90)~~.
8. **(Currently amended)** The isolated polypeptide of claim 6 comprising the amino acid sequence of the polypeptide of SEQ ID NO:90 ~~shown in FIG. 90 (SEQ ID NO:90)~~, lacking its associated signal peptide.
9. **(Currently amended)** The isolated polypeptide of claim 6 comprising the amino acid sequence of the extracellular domain of the polypeptide of SEQ ID NO:90, wherein the extracellular domain is amino acids 29-50 or 125-140 ~~shown in FIG. 90 (SEQ ID NO:90)~~.
10. **(Currently amended)** The isolated polypeptide of claim 6 comprising the amino acid sequence of the extracellular domain of the polypeptide of SEQ ID NO:90, wherein the extracellular domain is amino acids 29-50 or 125-140 ~~shown in FIG. 90 (SEQ ID NO:90)~~, lacking its associated signal peptide.

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11. **(Previously presented)** The isolated polypeptide of claim 6 comprising the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 209399.

12. **(Previously presented)** A chimeric polypeptide comprising a polypeptide according to claim 1 fused to a heterologous polypeptide.

13. **(Currently amended)** The chimeric polypeptide of claim 12, wherein said heterologous polypeptide is an ~~epitope~~ tag polypeptide or an Fc region of an immunoglobulin.